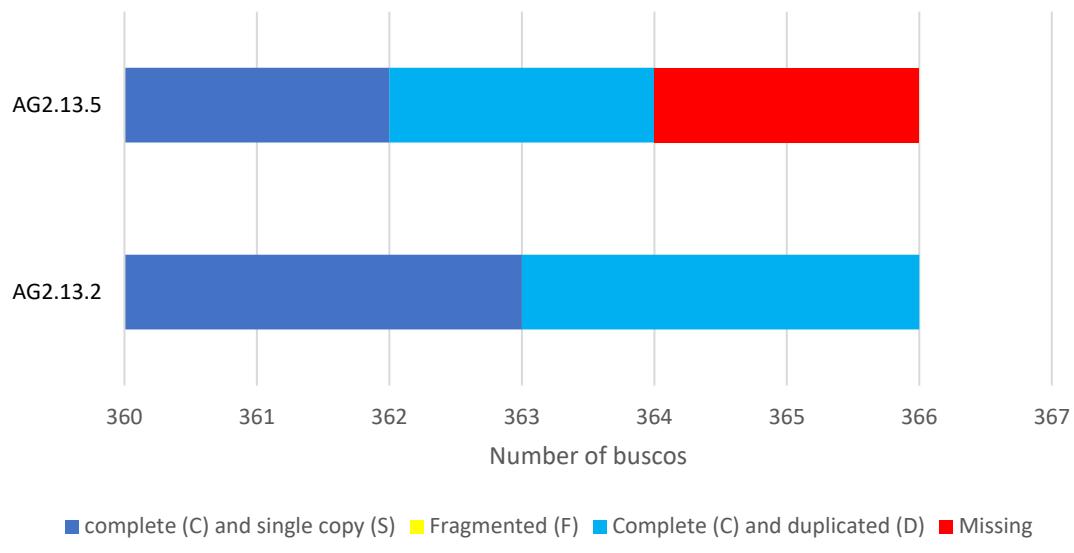


### Busco Assessment results



**Figure S1.** Busco Assessment results showing the number of copies of BUSCO genes complete ( $C = 366$ ), complete and single copy ( $S = 363$ ) complete and duplicated ( $D = 3$ ), fragmented ( $F=0$ ) and missing ( $M =0$ ) ( $n = 366$ ) in strain AG2.13.2 and complete ( $C = 364$ ), complete and single copy ( $S= 362$ ) complete and duplicated ( $D = 2$ ), fragmented ( $F=0$ ) and missing ( $M =2$ ) ( $n = 366$ ) in strain AG2.13.5.

**Table S1.** List of genomes downloaded from the NCBI- Characteristics of the strains selected for genome similarity assessments and comparisons.

Strain	GeneBank Assembly Accession number	Observations	Host/isolation source	Location	Date of isolation
<i>A. salmonicida</i> A527	GCA_002764135.1	Mesophilic	<i>Macrobrachium rosenbergii</i>	Mumbai, India	May 2007
<i>A. salmonicida</i> Y47	GCA_001481535.2	Mesophilic	<i>Gallus domesticus</i>	Mumbai, India	January 2006 to March 2008
<i>A. salmonicida</i> Y567	GCA_001466435.1	Mesophilic	Fish	Mumbai, India	January 2006 to March 2008
<i>A. salmonicida</i> Y577	GCA_001481545.2	Mesophilic	Fish	Mumbai, India	January 2006 to March 2008
<i>A. salmonicida</i> pectinolytica 34mel	GCA_000447435.2	Mesophilic, pigment producer	Heavily polluted water, Matanza river	Buenos Aires, Argentina	1988
<i>A. salmonicida</i> SWR-OG1	GCA_012931585.1	Mesophilic	<i>Epinephelus coioides</i>	Southern China Zhangzhou city, Fujian province	April 2018
<i>A. salmonicida</i> O23A	GCA_002180335.1	Dissimilatory arsenate reductase gene	Rock biofilm and bottom sediments	Zloty Stok gold mine in south-west Poland	2010
<i>A. salmonicida</i> S121	GCA_002214245.1	Unavailable. Most likely belonging to <i>salmonicida</i> subspecies	<i>Salmo salar</i>	Yantai, China	August 2015
<i>A. salmonicida</i> masoucida RZ6S-1	GCA_019443825.1	Intermediate	<i>Scophthalmus maximus</i>	Yantai, China	December 2016
<i>A. salmonicida</i> salmonicida J409	GCA_009858115.1	Unavailable	<i>Anoplopoma fimbria</i>	British Columbia, Vancouver, Canadá	2018

<i>A. salmonicida</i> <i>salmonicida A449</i>	GCA_000196395.1	Psychrophilic	<i>Salmo trutta</i>	Eure river, France	1975
<i>A. salmonicida</i> <i>salmonicida SHY16-3432</i>	GCA_008370735.1	Psychrophilic	<i>Salvelinus fontinalis</i>	Quebec, Canadá	2016
<i>A. salmonicida AS1</i>	GCA_029153695.1	Psychrophilic	Duck meat	Shandong, China	2016
<i>A. media T0.1-19</i>	<u>GCA_013085485.1</u>	Misidentified	Sludge from bioreactor treating oxytetracycline bearing wastewater	Beijing, China	April 2016
<i>A. media T5-1</i>	<u>GCA_019455365.1</u>	Misidentified	Biofilm of synthetic bioreactor under oxytetracycline stress	Beijing, China	May 2017
<i>A. media E31</i>	<u>GCA_016653695.1</u>		Water	Jinan, China	May 2018
<i>A. media WS</i>	<u>GCA_000287215.3</u>	High levels of melanin	Water	East Lake, Wuhan, Hubei, China	2003
<i>A. media R25-3</i>	<u>GCA_013085765.1</u>	Synthetic bioreactors under oxytetracycline stress	Sludge from bioreactor treating oxytetracycline bearing wastewater	Beijing, China	October 2017
<i>A. media R1-18</i>	<u>GCA_013085725.1</u>	Synthetic bioreactors under oxytetracycline stress	Sludge from bioreactor treating oxytetracycline bearing wastewater	Beijing, China	May 2016
<i>A. media R1-26</i>	<u>GCA_014109865.1</u>	Synthetic bioreactors under oxytetracycline stress	Biofilm reactor	China	July 2018
<i>Aeromonas rivipollensis KN-Mc-11NI</i>	GCA_003015165.1		<i>Myocastor coypus</i>	Nakdong River, South Korea	July 2017
<i>Aeromonas rivipollensis G42</i>	GCA_010974915.1		Water	Johannesburg, South Africa	March 2018
<i>Aeromonas veronii</i> bv <i>sobria</i> AG5.28.6	GCA_003367135.1		<i>Dicentrarchus labrax</i>	East Aegean Sea, Greece	November 2015
<i>Aeromonas veronii</i> bv <i>sobria</i> BIOO050A	GCA_004379215.1		<i>Dicentrarchus labrax</i>	East Aegean Sea, Greece	2009
<i>Aeromonas veronii</i> bv <i>sobria</i> NS6.15.2	GCA_004379225.1		<i>Dicentrarchus labrax</i>	West Aegean Sea, Greece	November 2015
<i>Aeromonas veronii</i> bv <i>sobria</i> NS	GCA_003367145.1		<i>Dicentrarchus labrax</i>	West Aegean Sea, Greece	2009
<i>Aeromonas veronii</i> bv <i>sobria</i> NS2	GCA_004379205.1		<i>Dicentrarchus labrax</i>	West Aegean Sea, Greece	September 2015
<i>Aeromonas veronii</i> bv <i>sobria</i> NS13	GCA_008119755.1		<i>Dicentrarchus labrax</i>	West Aegean Sea, Greece	2015
<i>Aeromonas veronii</i> bv <i>sobria</i> NS22	GCA_008119745.1		<i>Dicentrarchus labrax</i>	West Aegean Sea, Greece	September 2016
<i>Aeromonas veronii</i> bv <i>sobria</i> PDB	GCA_003367105.1		<i>Dicentrarchus labrax</i>	West Aegean Sea, Greece	2009
<i>Aeromonas veronii</i> bv <i>sobria</i> VCK1	GCA_003367095.1		<i>Dicentrarchus labrax</i>	East Aegean Sea, Greece	September 2015

Table S2. Statistics of the genome-assembly report.

Strain	Platforms	Assembler	Contigs	Largest Contig (bp)	N50	GC%	coverage	Number of reads
<i>A. salmonicida</i> AG2.13.2	Illumina	Unicycler v0.4.8	65	402.057	168.750	58.74	256.397	6214779

<i>A. rivipollensis</i> AG2.13.5	Illumina	Unicycler v0.4.8	50	501.196	288.897	61.61	243.19	5502017
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**Table S3.** *A. salmonicida* AG2.13.2 CRISPR site prediction.

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CRISPR ID	Start	End	Length(bp)	Number of spacers	Direct repeat length(bp)
<i>CRISPR2</i>	396098	396417	319	5	23

**Table S4.** *A. salmonicida* AG2.13.2 PHASTER prediction.

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PHASTER region	Start	End	Length(bp)	Number of Phage Hit Proteins
<i>INCOMPLETE PROPHAGE</i>	4627372	4669096	41,700 pb	41

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